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SEQUENCE LISTING

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<120> EPITHELIAL CELL GROWTH INHIBITORS

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<222> (774)

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<221> UNSURE

<222> (783)

<223> Xaa is any amino acid or may be absent

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Ile	Arg	His	Glu	His	Gly	Glu	Glu	Thr	Xaa	Glu	Val	Xaa	Asn	Lys	Xaa
1				5					10					15	

Glu	Ala	Pro	Gly	Ala	Pro	Pro	Val	Ser	Pro	Arg	Gly	Ala	Arg	Gly	Gly
			20						25					30	

Xaa	Arg	Arg	Pro	Cys	Gly	Pro	Pro	Val	Lys	Tyr	His	Tyr	Ser	Asp	Arg
			35					40						45	

Xaa	Xaa	Thr	Asp	Pro	Val	Arg	Arg	Gly	Gly	Glu	Pro	Arg	Gly	Ala	Leu
		50					55					60			

Ala	Xaa	Gly	Ala	Lys	Arg	Pro	Ala	Ala	Arg	Arg	Pro	Gly	Ala	Thr	Arg
	65					70				75					80

Ser	Gly	Xaa	Ser	Ala	Arg	Trp	Gly	Val	Xaa	Leu	Gly	Arg	Tyr	Thr	Cys
						85				90					95

Gln	Thr	Xaa	Xaa	Gln	Val	Ser	Xaa	Gly	Glu	Leu	Arg	Glu	Asp	Arg	Asn
				100					105						110

Leu Pro Trp Xaa Arg Arg Ala Lys Ala Arg Leu Ile Leu Ile Phe Ser
 115 120 125

Thr Asn Thr Asp Xaa Glu Ser Gly Ala Ser Arg Ser Phe Xaa Pro Phe
 130 135 140

Gly Phe Xaa Ala Xaa Xaa Val Arg Lys Val Thr Thr Gly Ile Thr Gly
 145 150 155 160

Leu Trp Arg Pro Ser Xaa His Ser Asp Val Ala Phe Xaa Ser Phe Asp
 165 170 175

Val Gly Ser Ser Tyr His Xaa Glu Ala Glu Phe Thr Lys Arg Trp Ile
 180 185 190

Val His Pro Leu Ile Gly Xaa Xaa Ser Trp Val Xaa Thr Val Val Arg
 195 200 205

Gln Val Ser Phe Thr Leu Leu Xaa Met Cys Cys Cys His Gly Asn Pro
 210 215 220

Ala Gln Tyr Glu Arg Asn Arg Arg Xaa Arg His Leu Val Tyr Val Leu
 225 230 235 240

Gly Xaa Gly Ala Asn Gly Ala Lys Xaa Xaa Ser Val Gly Leu Xaa Leu
 245 250 255

Asn Ala Ser Lys Ser Glu Ser Arg Pro Xaa Gly Thr Ile Arg Gln Arg
 260 265 270

Arg Gly Ala Ser Val Gly Leu Gly Xaa Pro Xaa Pro Arg Leu Ser Pro
 275 280 285

Pro Ala Gly Arg Pro Pro Pro Ser Thr Arg Xaa Xaa Arg Ala Gly Gly
 290 295 300

Arg Val Pro Arg Arg Ala Pro Gly Pro Gly Ser Xaa Ala Glu Cys Pro
 305 310 315 320

Ser Ser Trp Glu Thr Gly Arg Gly Arg Lys Gly Gly Xaa Pro Leu Ala
 325 330 335

Arg His Ala Pro His Val Arg Ala Arg Ala Glu Phe Xaa Xaa Ser Ser
 340 345 350

Thr Ile His Asn Arg His Thr Ser Ala Cys Ile Phe Met Xaa Xaa Xaa
 355 360 365

Ile	Leu	Phe	Leu	Trp	Val	Asp	Ile	Gln	Xaa	Trp	Asp	Cys	Xaa	Xaa	Thr	370	375	380	
Trp	Xaa	Phe	Tyr	Phe	Trp	Phe	Ile	Glu	Lys	Ser	Ser	Tyr	Xaa	Xaa	Xaa	385	390	395	400
Xaa	Arg	Leu	Tyr	Lys	Phe	Thr	Ser	Leu	Pro	Ser	Asp	Phe	Phe	Lys	Xaa	405	410	415	
Glu	Arg	Met	Val	Trp	Arg	Asn	Ala	Pro	His	Xaa	Tyr	Pro	Pro	Phe	Thr	420	425	430	
Xaa	Leu	Leu	Gln	Asn	Asp	Phe	Lys	Gly	Tyr	Arg	Tyr	Leu	Gln	Val	Ser	435	440	445	
Xaa	Xaa	Arg	Gln	Ile	Glu	Tyr	Xaa	Asn	Phe	Cys	Ile	Arg	Gly	Thr	Asp	450	455	460	
Phe	Xaa	Ile	Gln	Ser	Cys	Met	Asn	Lys	Asp	Lys	Cys	Ser	Arg	Asp	Leu	465	470	475	480
Gln	Ser	Xaa	Asn	Trp	Lys	Ser	Gln	Met	Lys	Tyr	Ile	Ser	Ser	Ser	Thr	485	490	495	
Thr	Ser	Xaa	Xaa	Ser	Thr	Glu	Leu	Ala	Leu	Xaa	Ser	Ser	Leu	Ile	Pro	500	505	510	
Thr	Tyr	Xaa	Xaa	Xaa	Xaa	Lys	Gly	Phe	Ile	Ser	Asn	Ile	Leu	Xaa	Gly	515	520	525	
Ile	Lys	Ile	Lys	Xaa	Xaa	Val	Lys	Leu	Phe	Ser	Leu	Ala	Phe	Xaa	Phe	530	535	540	
Gln	Asn	Ile	Lys	Xaa	Xaa	Pro	Ser	Ile	Gly	His	Leu	Tyr	Cys	Thr	Arg	545	550	555	560
His	Cys	Val	Cys	His	Xaa	Ser	Lys	Met	Phe	Ser	Trp	Xaa	Cys	Ser	Gln	565	570	575	
Xaa	Phe	Cys	Arg	Val	Arg	Xaa	Ser	Leu	Thr	Val	Val	Arg	Leu	Phe	Ser	580	585	590	
Lys	Arg	Asn	Leu	Xaa	Thr	Xaa	Xaa	Phe	Asn	Leu	Arg	Lys	Val	Ser	Asn	595	600	605	
Arg	Thr	Arg	Thr	Xaa	Thr	Xaa	Xaa	Ile	Thr	Leu	Gln	Ile	Ser	Pro	Tyr	610	615	620	

His Thr Ala Ser Thr Cys Ala Cys Xaa Leu Ile Pro Gly Ser Cys Tyr
625 630 635 640

Phe Pro Phe Tyr Phe Leu Ser Leu Xaa Xaa Thr Thr Pro Phe Ser Pro
645 650 655

His Phe Phe Ser Phe Phe Leu Ile Val Xaa Tyr Ile Thr Asn Thr Cys
660 665 670

Leu Ser Glu Gln Leu Ile Xaa His Lys Arg Xaa Xaa Ser Thr Gly Glu
675 680 685

Xaa Xaa Leu Ile Pro Val Ile Leu Ala Leu Xaa Xaa Ala Lys Ala Gly
690 695 700

Arg Ser Leu Glu Ser Arg Val Arg Asp Gln Pro Xaa Gln His Gly Glu
705 710 715 720

Thr Leu Ser Leu Gln Lys Asn Thr Lys Ile Xaa Pro Xaa Val Leu Ala
725 730 735

His Thr Cys Ser Leu Ser Tyr Ser Glu Gly Xaa Gly Xaa Xaa Ile Asp
740 745 750

Xaa Ala Gln Glu Val Glu Ala Ala Ala Val Arg Xaa Asp Xaa Ala Ile
755 760 765

Ala Leu Gln Pro Gly Xaa Glu Arg Glu Thr Leu Ser Gln Lys Xaa Lys
770 775 780